FIG. 6A

FIG. 6B

F16.6C

Centriolin Nud1-GFP-expressing cells



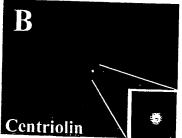




FIG. 6D

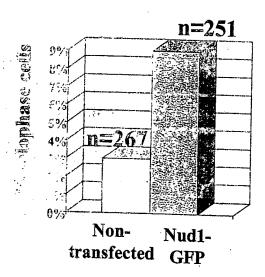
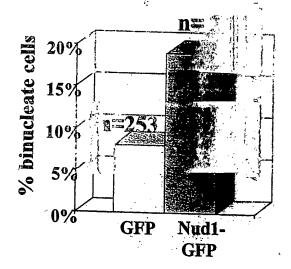
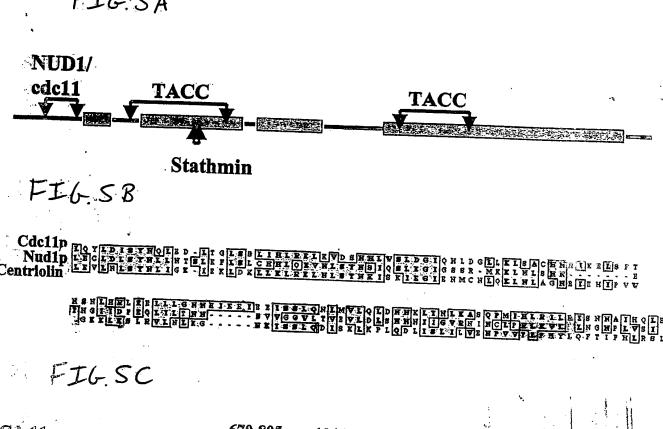


FIG. 6E



FREE !



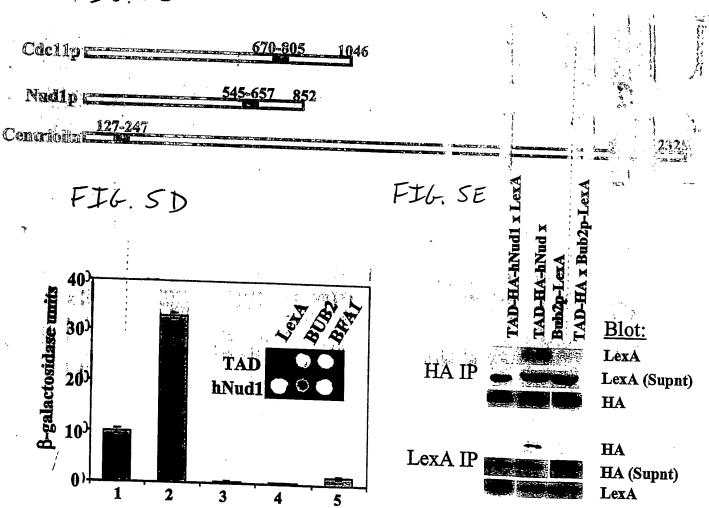
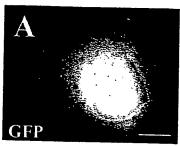


FIG. 6A

FIG. 6B

FI6.6C

Centriolin Nud1-GFP-expressing cells





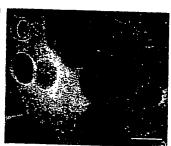
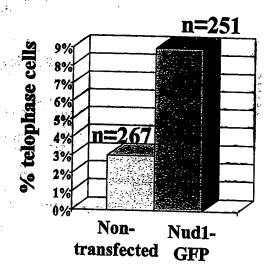


FIG. 6D



FJG. 6E

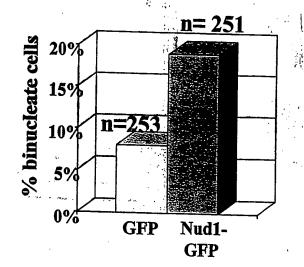
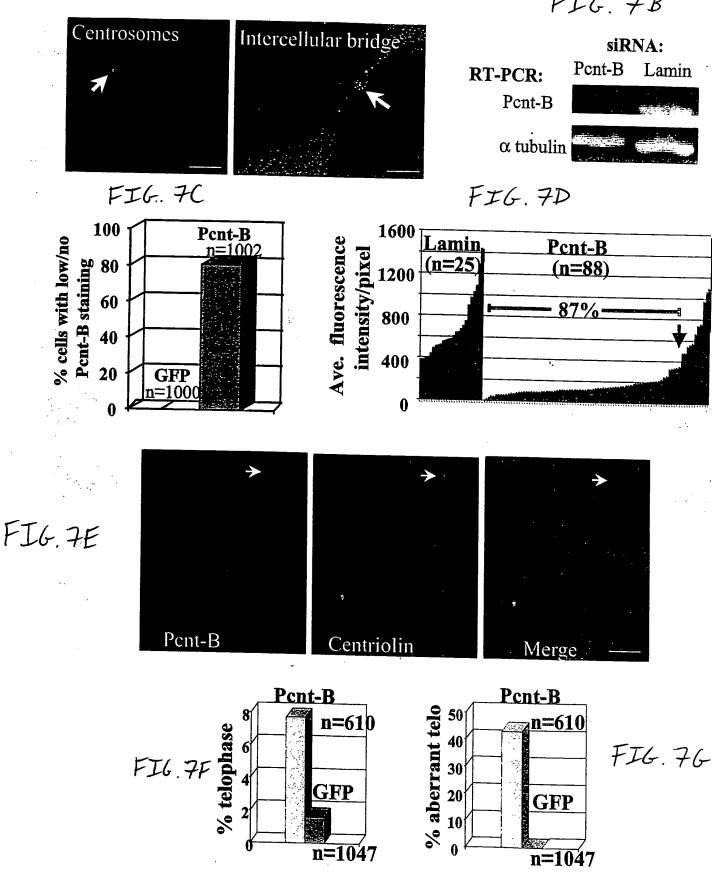
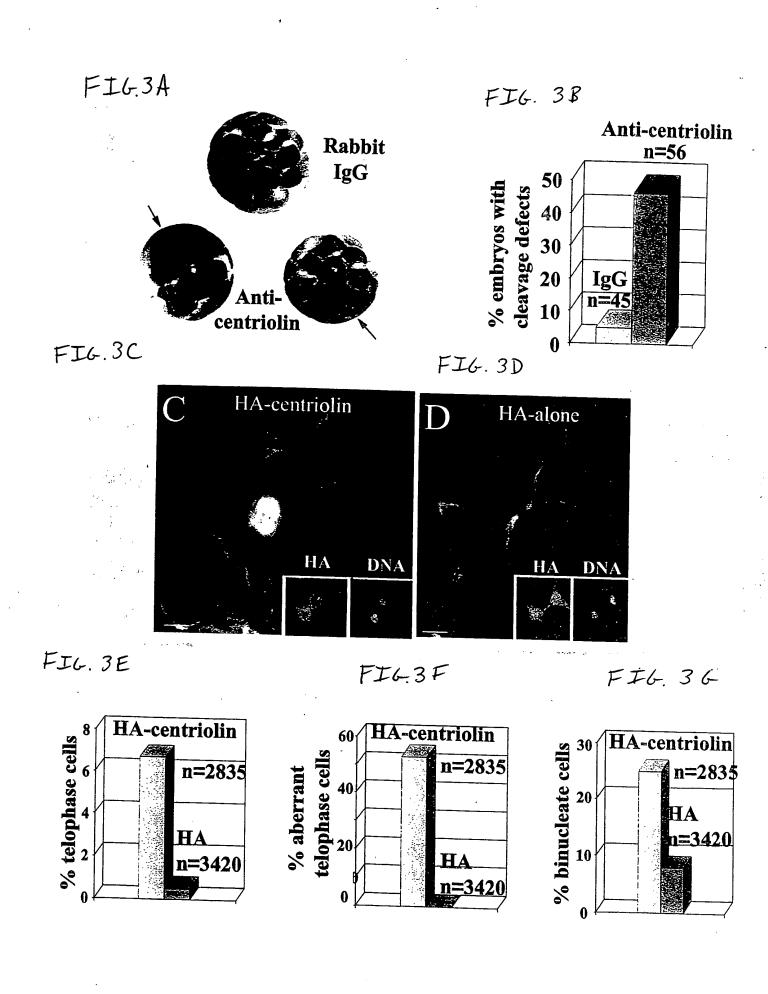


FIG. 7B

n=1047



n=1047



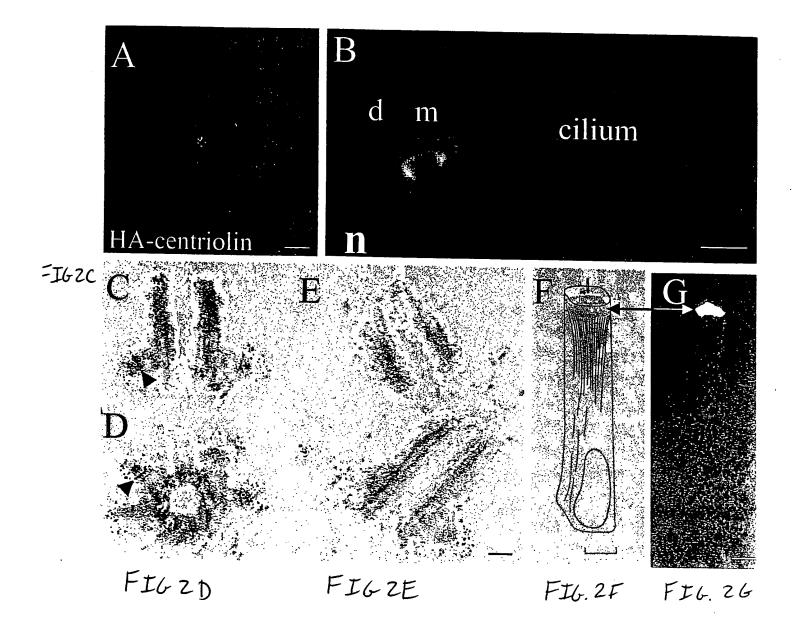
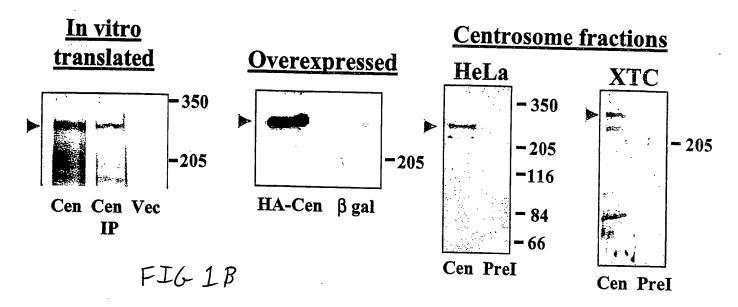
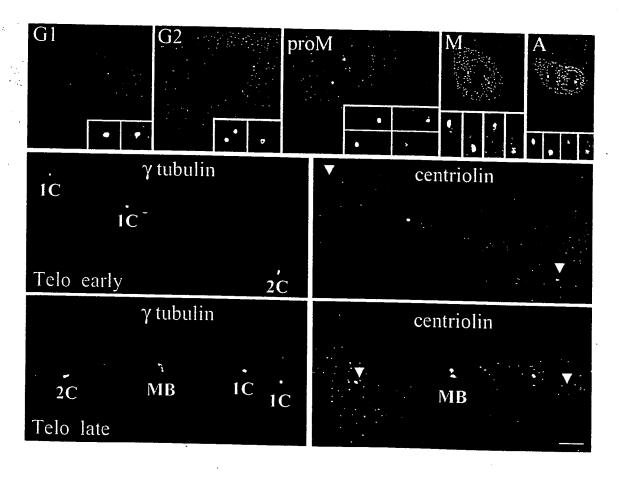
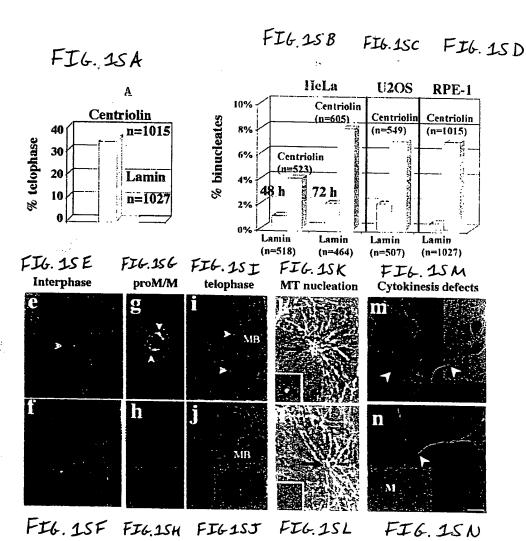
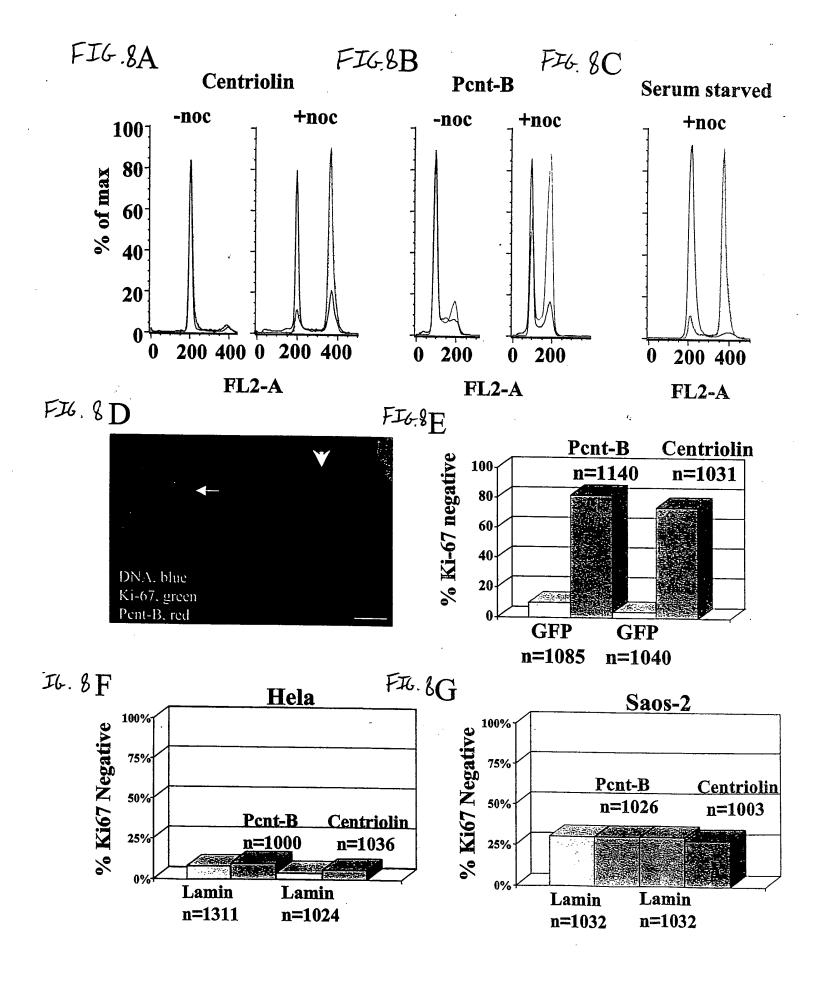


FIG 1A









Without nocodazole

	% G1	% S	% G2	
Centriolin	74	13	7	
Lamin (control)	62	23		
	% G1	% S	% G2	
Pent-B	70	13	14 19	
GFP (control)	46	35		

With nocodazole

	% G1	% S	% G2	
Centriolin	49	13	23	
Lamin (control)	19	19	46	
	% G1	% S	% G2	
Pent-B	65	14	19 60	
GFP (control)	24	15		

Cell cycle analysis of flow cytometry data (Fig 8).

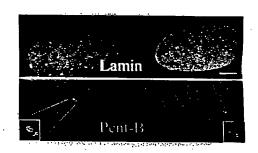


Figure 10

% cells with unaltered staining levels

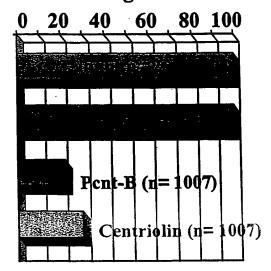


Figure 11

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F H S G G Q W C E Q I E I A D
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226
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R V L N L K G N K I S S L Q D
631
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676
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Ω11
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20.00

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 1261
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6166
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6211
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6301
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FIG. 12I

6706 gcactccgggagaaactgcgtcaccgggaagaccgactcaaggcc ALREKLRHREDRLKA 6751 Q L R H C M S K Q A E V L I K ggaaagcggcagacagagggcactttacacagtttgaggagacaa G K R Q T E G T L H S L R R Q 6841 $\verb|gtagatgctttaggggaattggtcaccagcacctctgcagattca|\\$ V D A L G E L V T S T S A D S 6886 gcgtcatcacccagtctgtctcagctggagtcttccctcacagag ASSPSLSQLESSLTE 6931 gactctcaacttggacaaaatcaggaaaagaatgcctcagccaga (SEQ ID NO:1) DSQLGQNQEKNASAR (SEQID NO:2) 6976 tga 6978

FIG. 13 A

1 atggaagttgagcaagagcagcggcagaaaggtggaggccggg $\begin{picture}(20,10) \put(0,0){\line(1,0){10}} \put(0$ 46 aggacgaagcttgctcacttccgacagagaaaacaaaaggtgac RTKLAHFRQRKTKGD 91 agttcgcattcggagaaaaagacggcgaagaggaagggctcggct SSHSEKKTAKRKGSA 136 gtcgatgcgtctgtccaggaggagagtccggtaaccaaggaggac V D A S V Q E E S P V T K E D 181 agcgcactctgtggaggaggggacatttgcaaaagcacatcatgt S A L C G G G D I C K S T S C 226 gacgacacccctgatggggcaggagggcctttgcagctcagccg D D T P D G A G G A F A A Q P 271 gaggactgtgatggagagagagagaggacttggaacagctgcag EDCDGEKREDLEQLQ 316 cagaagcaagtcaatgaccatectccagagcagtgtgggatgttc Q K Q V N D H P P E Q C G M F 361 acagtcagtgaccaccaccagaacagcatgggatgttcacagtc TVSDHPPEQHGMFTV GDHPPEQRGMFTVSD 451 cacccaccagaacagcatgggatgttcacagtcagtgaccaccca H P P E Q H G M F T V S D H P 496 ccagaacagcgtgggatgttcacaatcagtgaccaccaaccggaa PEQRGMFTISDHQPE 541 cagcgtgggatgttcacagtcagtgaccacacaccagaacagcgt $\hbox{Q} \hbox{ R} \hbox{ G} \hbox{ M} \hbox{ F} \hbox{ T} \hbox{ V} \hbox{ S} \hbox{ D} \hbox{ H} \hbox{ T} \hbox{ P} \hbox{ E} \hbox{ Q} \hbox{ R}$ 586 gggatetteacaateagtgaceacecageagaacagegtgggatg $\hbox{\tt G} \hbox{\tt I} \hbox{\tt F} \hbox{\tt T} \hbox{\tt I} \hbox{\tt S} \hbox{\tt D} \hbox{\tt H} \hbox{\tt P} \hbox{\tt A} \hbox{\tt E} \hbox{\tt Q} \hbox{\tt R} \hbox{\tt G} \hbox{\tt M}$.631 ttcacaaaggagtgtgaacaagaatgtgaacttgccattactgac FTKECEQECELAITD 676 ctggagagcggccgtgaagatgaggctggcctgcatcagagtcag L E S G R E D E A G L H Q S Q 721 gccgtgcatggccttgagctggaggcgctgcgcctgagtctgagc A V H G L E L E A L R L S L S 766 aacatgcacacggcgcagctggagctgacacaggccaacctccag N M H T A Q L E L T Q A N L Q 811 aaggagaaggagacggcattgacggagctgcgggagatgctcaac S R R A Q E L A L L Q S R Q Q 901 cacgagetggageteeteagggageageaegeaeggagaaggag HELELLREQHAREKE E V V L R C G Q E A A E L K E 991 aagttacaatcagaaatggagaaaaacgcccagatagtaaagacc KLQSEMEKNAQIVKT 1036 ctgaaggaagattgggaatctgaaaaagatttatgtttagaaaat L K E D W E S E K D L C L E N 1081 ctacgcaaagaactgtctgcaaagcatcaatcagaaatggaggat LRKELSAKHQSEMED 1126 ttacaaaaccagtttcagaaagaattggcagaacagagagctgag LQNQFQKELAEQRAE 1171 ttggagaagatttttcaagacaaaaaccaggctgaacgggccctt LEKIFQDKNQAERAL 1216 aggaacctggagagtcatcatcaagcagccattgagaagttacgt RNLESHHQAAIEKLR

1261 gaagacctgcagtccgagcacggccggtgtttagaagacttggag E D L Q S E H G R C L E D L E 1306 ttcaagttcaaagagagcgagaaagaaaaacagctggagttagag F K F K E S E K E K Q L E L E 1351 aatcttcaagcatcatatgaagacctgaaggcacaatcacaagaa N L Q A S Y E D L K A Q S Q E 1396 gagatcaggcgcttgtggtcccagcttgattctgccaggaccagt E I R R L W S Q L D S A, R T S 1441 agacaggaattgagtgagctacatgagcaactcctggcgcgcacc RQELSELHEQLL'ART 1486 tctcgtgtggaagatttagaacagctgaagcagcgagaaaaaacc SRVEDLEQLKQREKT 1531 cagcatgagtccgaactggagcaactgaggatttattttgaaaag Q H E S E L E Q L R I Y F E K 1576 aagttaagggatgctgagaaaacttaccaagaagacctaaccctg KLRDAEKTYQEDLTL 1621 ttacagcagaggctgcagggggggggaagatgctcttctggac LQQRLQGAREDALLD 1666 tctgtggaagttgggttgtcctgtgtgggtttagaagagaaacct SVEVGLSCVGLEEKP 1711 gagaaaggaagaaaagatcacgttgatgaactcgagcctgagcga EKGRKDHVDELEPER 1756 cataaggagagcctgccacgcttccaggcggagttagaagaagc H K E S L P R F Q A E L E E S 1801 cacaggcaccagctggaagcgctggagtctcccctctgcatccag H R H Q L E A L E S P L C I Q 1846 cacgagggcatgtctcagacagatgctgcgtagagacttcagca 1891 ttgggacacgagtggcgtctggaaccctctgaagggcacagccaa L G H E W R L E P S E G H S Q 1936 gagetteeetgggtgcateteeagggtgtgcaggacggggaettg E L P W V H L Q G V Q D G D L 1981 gaggccgacacagagcgggcagccagagtcttgggtctggaaact EADTERAARVLGLET 2026 gagcacaaggtgcaactttcgcttcttcagactgagctcaaagaa E H K V Q L S L L Q T E L K E 2071 gaaattgaactcctaaaaatagaaaatagaaatttgtatgagaag EIELLKIENRNLYEK 2116 ttgcagcatgaaactcgtctgaaggacgatttggagaaggtaaaa LQHETRLKDDLEKVK 2161 cacaatctaattgaagaccaccagaaggaactaaataatgctaag H N L I E D H Q K E L N N A K 2206 caaaagactgagctgatgaaacaggaattccaaagaaaagaaacg QKTELMKQEFQRKET 2251 gactggaaagttatgaaggaggagctacagcgggaagctgaggag DWKVMKEELQREAEE 2296 aagttaacattgatgctacttgaactgagagaaaaggctgaatcc K L T L M L L E L R E K A E S 2341 gagaaacagaccatcataaacaagtttgagcttcgagaagctgaa EKQTIINKFELREAE 2386 atgaggcagcttcaggaccaacaggcagcccagatcctggatctg MRQLQDQQAAQILDL 2431 gagaggteettgaeggageageaggeegeetgeageagetggaa E R S L T E Q Q G R L Q Q L E 2476 caggacctcacttcagacgacgccctgcattgcagccagtgtggg Q D L T S D D A L H C S Q C G

2521 cgggagccgcccacagcccaggacggggagcttgccgcgctccac R E P P T A Q D G E L A A L H 2566 gtgaaggaagactgcgccctgcagctgatgctggcccggagcagg V K E D C A L Q L M L A R S R 2611 tttttagaggaacgtaaagagatcaccgagaaattcagtgcggaa FLEERKEITEKFSAE 2656 caagatgccttcctgcaggaggcccaggagcagcatgcccgtgag Q D A F L Q E A Q E Q H A R E 2701 ctgcagctcctccaggagagacaccagcagcagctcctgtcagtg L Q L L Q E R H Q Q Q L L S V 2746 acggcggagctcgaggccagacaccaggccgcgttgggcgagctg TAELEARHQAALGEL 2791 acagectecttagagageaageagggggetetgetggetgeaegt TASLESKQGALLAAR 2836 gtggccgaactgcagacaaaacacgctgccgacctcggcgctctg 2881 gagaccagacatctgtccagccttgattctttggaatcctgttac E T R H L S S L D S L E S C Y LSEFQTIREEHRQAL 2971 gagctcttacgagcagactttgaggaacaactgtggaaaaaggac ELLRADFEEQLWKKD 3016 tctcttcaccaaacgattttgactcaagagttggagaaactgaag SLHQTILTQELEKLK 3061 cggaaacacgaaggggagctacagtctgtgcgggaccacctgcga R K H E G E L Q S V R D H L R 3106 accgaagtgagcacagagctcgccggaaccgtggctcacgagctg TEVSTELAGTVAHEL 3151 cagggagtgcaccagggtgaatttggaagtgaaaagaaactgct Q G V H Q G E F G S E K K T A 3196 ttgcatgaaaaagaggagacacttcggcttcagagtgcacaggca LHEKEETLRLQSAQA 3241 cagccttttcaccaagaggagaaagagtctttgtctctgcagctt Q P F H Q E E K E S L S L Q L 3286 caaaagaagaatcaccaagtccagcagctgaaagaccaggtttta $\begin{smallmatrix} \mathsf{O} & \mathsf{K} & \mathsf{K} & \mathsf{N} & \mathsf{H} & \mathsf{O} & \mathsf{\Lambda} & \mathsf{O} & \mathsf{O} & \mathsf{\Gamma} & \mathsf{K} & \mathsf{D} & \mathsf{O} & \mathsf{\Lambda} & \mathsf{\Gamma} \end{smallmatrix}$ 3331 tccttaagtcacgagatagaagagtgccgctccgagttggaggtg SLSHEIEECRSELEV 3376 ctgcagcagaggcgggagcgggagaaccgggaaggcgcaaacctc L Q Q R R E R E N R E G A N L 3421 ctctccatgctcaaggccgacgtcaacctgtcccacagcgaaaga LSMLKADVNLSHSER G A L Q D A L R R L L G L F G 3511 gagacgctgagggcagccgtcaccctgaggagccggatcggggag $\hbox{\tt E} \quad \hbox{\tt T} \quad \hbox{\tt L} \quad \hbox{\tt R} \quad \hbox{\tt A} \quad \hbox{\tt A} \quad \hbox{\tt V} \quad \hbox{\tt T} \quad \hbox{\tt L} \quad \hbox{\tt R} \quad \hbox{\tt S} \quad \hbox{\tt R} \quad \hbox{\tt I} \quad \hbox{\tt G} \quad \hbox{\tt E}$ 3556 cgcgtggggctctgcctggatgacgcgggcgcaggcctggccctg R V G L C L D D A G A G L A L 3601 tcgacagctccggcgctggaggagacatggtctgatgtggccctc S T A P A L E E T W S D V A L 3646 ccggagttggacagaactttgtctgaatgtgcagagatgtcttcc PELDRTLSECAEMSS 3691 gtggctgaaattagcagccacatgcgtgaaagctttctcatgagc V A E I S S H M R E S F L M S 3736 ccagaaagtgtgcgggagtgtgagcagcccatccggagggtcttc PESVRECEQPIRRVF

3781 cagagectcagectggccgtggacggcctcatggagatggccctg Q S L S L A V D G L M E M A L D S S R Q L E E A R Q I_| H S R 3871 tttgaaaaagaatttagttttaagaatgaggagacagcacaggtt F E K E F S F K N E E T A Q V 3916 gtcaggaagcaccaggagctgctggagtgtttgaaggaggagagc V R K H Q E L L E C L K E E S 3961 gcagcaaaggcagagctggcgctggagctgcacaagactcagggt A A K A E L A L E L H K T Q G 4006 acccttgagggattcaaggtggagacagcagatctgaaggaggtg TLEGFKVETADLKEV 4051 ctggccgggaaggaggattccgagcaccgtctggtgctggagctg LAGKEDSEHRLVLEL 4096 gagagectgagacggeagetgeageaggeggeecaggageaggeg ESLRRQLQQAAQEQA ALREECTRLWSRGEA TATDAEAREAALRKE 4231 gtggaggatctgaccaaagaacagtcggagaccaggaagcaggct V E D L T K E Q S E T R K Q A 4276 gagaaggaccgctcagccctgctctcccagatgaagattttggag EKDRSALLSQMKILE 4321 tctgagttagaagaacagctgtctcagcatcgcgggtgtgccaag S E L E E Q L S Q H R G C A K 4366 caggcggaggccgtcactgccctggaacagcaggtggcatctctg Q A E A V T A L E Q Q V A S L 4411 gacaagcatttgcgcaaccagcggcaattcatggatgagcaggca D K H L R N Q R Q F M D E Q A 4456 gccgagcgggagcacgagcgcgaggagttccagcaggagattcag A E R E H E R E E F Q Q E I Q 4501 aggctggagggcagctccgccaggcggccaagccgcagccctgg RLEGQLRQAAKPQPW 4546 ggccctcgcgacagccagcaggcgccgctggatggagaggttgag G P R D S Q Q A P L D G E V E 4591 ttgttacaacaaaagttgagagaaaagttggatgaatttaatgaa LLQQKLREKLDEFNE 4636 ttggctatacagaaagagtcggcagatagacaagtgttaatgcag 4681 gaagaagaaattaaacgtctggaggagatgaacatcaacatcagg EEEIKRLEEMNINIR 4726 aaaaaagtggcccagctccaggaagaagtggaaaaacagaaaaac K K V A Q L Q E E V E K Q K N 4771 atcgtgaaagggctggaacaggataaagaggtgttaaagaaacag IVKGLEQDKEVLKKQ 4816 cagatgagtagettgettetggegteeaegttgeagtetaeaeta $\begin{smallmatrix} Q & \mathbb{M} & S & S & L & L & L & A & S & T & L & Q & S & T & L \\ \end{smallmatrix}$ 4861 gatgcaggcagatgtcccgagcctccttcgggcagccctcctgag D A G R C P E P P S G S P P E 4906 ggtccagaaatacagttagaggtgacacagagagcactcctgcgg G P E I Q L E V T Q R A L L R 4951 cgcgagagcgaggttttggacttaaaagaacagctagaaaagatg RESEVLDLKEQLEKM 4996 aaaggtgacttagaaagtaaaaatgaagaaatactacatctgaac KGDLESKNEEILHLN

5041 ttaaaattggacatgcagaacagccagactgctgtcagcctcaga 5086 gaacttgaggaagagaacacgagcttgaaggtcatatataccaga E L E E E N T S L K V I Y T R 5131 agttctgagattgaagagctgaaagccactattgaaaatctgcaa SSEIEELKATIENLQ 5176 gagaatcagaaacgattacaaaaggagaaagcagaggaaattgaa ENQKRLQKEKAEEIE 5221 caactccatgaagtcattgagaagctgcagcacgagctgtccctc Q L H E V I E K L Q H E L S L 5266 atggggcctgtggtgcacgaagtcagcgacagtcaggctggcagt MGPVVHEVSDSQAGS 5311 ctgcagagcgagctgctctgctcccaggccgggggccctcgtggg L Q S E L L C S Q A G G P R G 5356 caggccctacagggcgagctcgaggctgcgctggaagccaaggag QALQGELEAALEAKE 5401 gccctgagccggctgctggctgaccaggagcgcaggcacagccag A L S R L L A D Q E R R H S Q 5446 gccctggaggccctgcagcagcgcctccagggcgcagagggct A L E A L Q Q R L Q G A E E A 5491 gcggagctacagctggctgagctggagcgcaatgtagccctcagg A E L Q L A E L E R N V A L R 5536 gaggetgaggtegaagacatggeeteeeggateeaggagttegaa E A E V E D M A S R I Q E F E 5581 gcggccctgaaagcaaaggaagcgacgattgccgagagaaattta A A L K A K E A T I A E R N L 5626 gaaatcgacgctctgaaccagcggaaggcggcccactctgccgag E I D A L N Q R K A A H S A E 5671 ctggaggccgtcctgttggccttggcccgcatccgccgcgccctg LEAVLLALARIRRAL 5716 gagcagcagccctggcagccggggggggcgcctcccgagctgcag EQQPLAAGAAPPELQ 5761 tggctccgagcgcagtgtgcccgcctcagccgccagctgcaggtg WLRAQCARLSRQLQV 5806 ctgcaccagcggttcctgaggtgccaggtggagctggacaggcgg LHQRFLRCQVELDRR 5851 caggecegeagagecaeageteaeaegggtgeeeggggeeeae $\hbox{Q} \hbox{ A} \hbox{ R} \hbox{ R} \hbox{ A} \hbox{ T} \hbox{ A} \hbox{ H} \hbox{ T} \hbox{ R} \hbox{ V} \hbox{ P} \hbox{ G} \hbox{ A} \hbox{ H}$ 5896 ccacagcetegeatggatggtggccaaggcccaggtcaccggc 5941 gacgtggaggcctcccatgatgctgctttggagccggttgtccct 5986 gacccacagggtgatctgcagcctgtcctggtgacgttgaaggat 6031 gcacctctctgcaagcaagaaggcgtgatgtcagtgctcaccgtc 6076 tgccagaggcagctgcagtcggagctgctcttggtgaaaaatgaa CQRQLQSELLLVKNE 6121 atgcgcctgagtctggaggacggcggcaagggtaaagaaaagta 6166 ctggaagattgtcagctgccgaaggtcgatctcgtagctcaggtg L E D C Q L P K V D L V A Q V 6211 aaacagcttcaggaaaaactgaaccgtttgctgtattccatgacc KQLQEKLNRLLYSMT 6256 ttccagaatgtggatgctgccgacaccaaatctctgtggcccatg F Q N V D A A D T K S L W P M

6301 gcctcagcacacctgttggagagcagctggagtgatgattcctgt A S A H L L E S S W S D D S C 6346 gacggagaagagcctgacatatcaccccacatagacacatgtgat D G E E P D I S P H I D_i T C D 6391 qccaatacagccacggggggtgtaactgatgttatcaaaaatcag ANTATGGVTDVIKNQ 6436 gccatagacgcgtgtgatgccaatacaaccccagggggtgtaact AIDACDANTTPGGVT 6481 gatgttatcaaaaattgggattccttgataccagatgaaatgcca $\begin{picture}(100,10) \put(0,0){\line(1,0){10}} \put($ 6526 gattctcccattcaagaaaaatcagaatgtcaggacatgtctctt D S P I Q E K S E C Q D M S L 6571 tcttcaccgaccagcgtacttggtggctcccgccaccagagccac S S P T S V L G G S R H Q S H 6616 actgcagaggctgggccccggaagagcccggtcgggatgctggac TAEAGPRKSPVGMLD 6661 ctgtcttcctggagctcccctgaggtcctcaggaaggactggacc LSSWSSPEVLRKDWT 6706 ctggagccctggcccagcctcccgtgacaccccactcaggagcc L E P W P S L P V T P H S G A 6751 ctgagcctgtgcagtgccgacacatccctggggggacagggcggac L S L C S A D T S L G D R A D 6796 acctcgctgccacagacccaggggccggggctgctttgttcccca T S L P Q T Q G P G L L C S P 6841 ggcgtgtctgcagcagcgctggcactgcagtgggccgagtctccg G V S A A A L A L Q W A E S P 6886 ccggctgacgaccaccatgtgcagaggacggctgtggagaaagat P A D D H H V Q R T A V E K D 6931 gtcgaagattttatcacaacatcctttgattctcaagaaacatta V E D F I T T S F D S Q E T L 6976 agttcacctcctcctggattagaaggaaaagctgatagaagtgag S S P P P G L E G K A D R S E 7021 aaaagtgacggctcgggttttggagcaagactgagcccggggtca 7066 ggaggccctgaggctcaaactgctggtcctgtgacccctgcttcc G G P E A Q T A G P V T P A S 7111 atctctggaaggtttcagccgctgccggaagccatgaaggagaag I S G R F Q P L P E A M K E K 7156 gaagtgcgtccgaagcacgtgaaggctttactgcagatggtgcgt $\hbox{\tt E} \quad \hbox{\tt V} \quad \hbox{\tt R} \quad \hbox{\tt P} \quad \hbox{\tt K} \quad \hbox{\tt H} \quad \hbox{\tt V} \quad \hbox{\tt K} \quad \hbox{\tt A} \quad \hbox{\tt L} \quad \hbox{\tt L} \quad \hbox{\tt Q} \quad \hbox{\tt M} \quad \hbox{\tt V} \quad \hbox{\tt R}$ 7201 gacgagagccaccagatcctggcgctgtcagaaggccttgcaccc D E S H Q I L A L S E G L A P 7246 ccaagcggcgagccacacccacccggaaggaagacgagatacag PSGEPHPPRKEDEIQ 7291 gacatctcgctccatgggggaaagacgcaggaagtgcccaccgcg DISLHGGKTQEVPTA 7336 tgccccgattggagaggggaccttctgcaggttgtgcaagaggcc C P D W R G D L L Q V V Q E A 7381 tttgaaaaagagcaggagatgcagggggttgagctgcagccccga $F \quad E \quad K \quad E \quad Q \quad E \quad M \quad Q \quad G \quad V \quad E \quad L \quad Q \quad P \quad R$ 7426 ctcagtggctcagatctggggggtcacagctccctgctcgaaagg L S G S D L G G H S S L L E R 7471 ctggagaagatcatccgtgagcagggagacctgcaggaaaagtcc LEKIIREQGDLQEKS 7516 ctggagcatcttcgcttgccggaccggagcagcctgctgtccgag LEHLRLPDRSSLLSE

7561 atccaggcgctgcgtgcccagctgcgcatgacgcacctgcagaac IQALRAQLRMTHLQN 7606 caggagaagctgcagcacttgcgcacggcgctgacaagcgcagag QEKLQHLRTALTSAE ARGSQQEHQLRRQVE 7696 ctgctggcttataaagtagagcaggagaagtgcattgctggtgac LLAYKVEQEKCIAGD 7741 ttgcagaagacgctgagtgaagagcaagagaaggcaaacagcgtg L Q K T L S E E Q E K A N S V 7786 cagaageteetggeggeggageagactgtagtgegagatttgaag Q K L L A A E Q T V V R D L K 7831 tccgacctctgtgagagcaggcagaagagcgaacagctgtcccgg S D L C E S R Q K S E Q L S R 7876 tecetetgegaggtgeageaggaggteeteeagetgagateeatg $\verb|SLCEVQQEVLQLRSM|$ 7921 ctgagcagtaaggagaacgagctgaaggccgcgcttcaggagctg LSSKENELKAALQEL 7966 gagagtgagcaggggaagggggtgccctgcagagccagctggag E S E Q G K G R A L Q S Q L E 8011 gaggagcagctgcggcacctgcagagggagagccagagtgccaag E E Q L R H L Q R E S Q S A K 8056 gccctggaggagctgcgggcgtctttggagacacagcgtgctcag ALEELRASLETQRAQ 8101 agcagtcgactctgcgtggcactgaaacacgagcagacggccaag SSRLCVALKHEQTAK 8146 gacaacctgcagaaggagctgcgtatcgagcactcacgctgcgag DNLQKELRIEHSRCE 8191 gccttgctggctcaggagcggagccagctctctgagctccagaag ALLAQERSQLSELQK 8236 gaccttgcggctgagaagagccgcaccctggagctgtcagaggcc DLAAEKSRTLELSEA 8281 ttgcggcacgagcggctcctgaccgagcagctgagccagaggaca LRHERLLTEQLSQRT 8326 caggaggettgcgtgcaccaggacacacaggcccatcacgctctg Q E A C V H Q D T Q A H H A L 8371 ctgcagaagctgaaggaggagaagtcccgggtggtggacttgcaa L Q K L K E E K S R V V · D L Q 8416 gcgatgcttgaaaaggtgcagcagcaagccctgcattctcagcag A M L E K V Q Q Q A L H S Q Q 8461 cagcttgaggctgaggctcagaagcactgtgaggcgctcaggaga Q L E A E A Q K H C E A L R R 8506 gagaaggaggtaagtgccacactgaagtcgacggtggaagccctg EKEVSATLKSTVEAL H T Q K R E L R C S L E R E R 8596 gagaaaccagcgtggttgcaggcagaattagagcagtcacaccca E K P A W L Q A E L E Q S H P 8641 cggttgaaagagcaagaaggacgcaaggctgcgaggaggagcgcg RLKEQEGRKAARRSA 8686 gaggccaggcagagcccagcggctgcggagcagtggaggaagtgg EARQSPAAEQWRKW 8731 cagagagacaaggagaagctgcgagaattagaactgcagcgtcag QRDKEKLRELELQRQ 8776 cgtgacttgcataagatcaagcagcttcagcagacagtgagagac RDLHKIKQLQQTVRD

FIG. 13H

8821 ctggagtcgaaggacgaggtgcctggcagccgcctccacctaggt LESKDEVPGSRLHLG 8866 tctgcccgcagggctgccggctcggatgcggaccacctccgggaa S A R R A A G S D A D H_i L R E 8911 cagcagcgagagctggaggcgatgaggcagcggctgctctctgcc QQRELEAMRQRLLSA 8956 gcccggcttctcaccagcttcaccagccaggccgtggacaggaca A R L L T S F T S Q A V D R T 9001 gttaatgattggacgtcatccaatgagaaagcagtgatgtcttta V N D W T S S N E K A V M S L 9046 ctgcacacgttggaggagctgaagtctgacttgagcaggcccacc LHTLEELKSDLSRPT 9091 tootoccagaaaaaaaatggcagcagagctgcagttccagtttgtg S S Q K K M A A E L Q F Q F V 9136 gacgtcctgctgaaagacaatgtttccctcacaaaagcgctcagc D V L L K D N V S L T K A L S 9181 acggtgacccaggagaagctggagctgagcagagccgtgtctaag TVTQEKLELSRAVSK 9226 cttgagaagttgctgaagcaccatctgcagaagggctgcagccca L E K L L K H H L Q K G C S P 9271 agcaggtcggaaaggtctgcttggaagccagacgaaacggctcca S R S E R S A W K P D E T A P 9316 cagagttccctgaggcgcccagaccccggccggcttccaccagct Q S S L R R P D P G R L P P A 9361 gccagcgaggaagcacaccagcaatgtcaagatggaaaaattg A S E E A H T S N V K M E K L 9406 tacctgcattacttgagagcagagagctttagaaaagctctgatt YLHYLRAESFRKALI 9451 tatcaaaagaagtatcttttgctgttgattggtggattccaggat Y Q K K Y L L L L G G F Q D 9496 totgaacaagaaacactotocatgattgcccatttggggggtattt S E Q E T L S M I A H L G V F 9541 ccttccaaagcagaacggaaaatcacatctcgtcctttcaccagg PSKAERKITSRPFTR 9586 ttccgcacggccgtcagggtggtcattgcaatattaagattacgt F R T A V R V V I A I L R L R 9631 tttttggttaagaaatggcaagaagtagatcggaaaggagctctg F L V K K W Q E V D R K G A L 9676 gcacaaggcaaagcccctcgcccagggccccgagcacgacagccg A Q G K A P R P G P R A R Q P 9721 cagtctccacccagaaccagagagtcccccccaacccgggatgta Q S P P R T R E S P P T R D V 9766 ccctctggccacaccagggaccctgccagaggccgcagactggca P S G H T R D P A R G R R L A 9811 gcagcagcctccccacacagtgggggaagagccactccatccca A A A S P H S G G R A T P S P 9856 aattcaagattagaaagatccctgactgcttctcaagatccagaa N S R L E R S L T A S Q D P E 9901 cattccttgacagagtatattcaccatttagaagtgatccagcaa H S L T E Y I H H L E V I Q Q 9946 agattgggaggggtactaccagattctacttcaaagaaatcctgc RLGGVLPDSTSKKSC (SEQ ID NO:3) 9991 cacccgatgattaaacagtga 10011 (SEQ ID NO:4) H P M I K Q

A	Nud1p/ Cdc11p 35%/50%)	TAC(Stathmin (40%/51%)	TACC (20%/41%)	A CONTRACTOR OF THE CONTRACTOR	
	127 233	435	623	879- 913	1385	1658	2,325
Coiled	coils 265- 349	434	801	845 1106	1317		2214

Figure 14

FIG. 16A

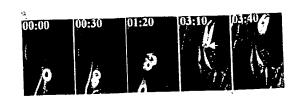
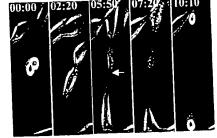
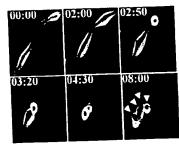


FIG. 26B



FI6.16C



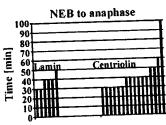


FIG. 26 E



FIG. 26 P

Metaphase to cleave Me 1400 500 500 400 300 Lamin 100 Time [min]

: = :-

FI6.16F

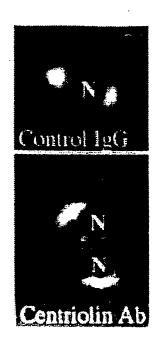


Figure 17